

10/573939
IAP20 Rec'd PCT/PTO 30 MAR 2006
Sequence protocol

<110> Epigenomics AG

<120> Method for the methylation analysis of DNA

<160> 5
<210> 1
<211> 20
<212> DNA
<213> Artificial Sequence

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<223> primer1

<400> 1

gggattattt ttataagggtt 20

<210> 2
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> primer2

<400> 2

tactaaaaac tctaaacccc atc 23

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<220>
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ttcgtcgctcg tagttttcggt t 21

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<220>
<223> detection probe2

<400> 4

tagtgagtagc gcgcgggtt 18

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<213> Artificial Sequence

<220>
<223> blocker

<400> 5

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25